54	GCC
	BCG
	TCT
45	CAG
	CCI
	GGG
36	GTG
	CCG
	CGC
27	
	GGT
	CCT
18	AGC
	CAA
(	$\mathcal{C}$
9	CAN
(	) 5
1414	NININ
. <u>-</u>	<b>1</b>

108 TCG S	162 CTC L	216 GTG V	270 ACT T	324 GAT D
9 9	CAC H	3 3 3	ATG M	TTG L
CCG	AAG K	ATT I	GTC V	
99 GCC A	153 CTG L	207 Gaa E	261 GAT D	315 TGG W
	GAG	ACA T	GAT D	AGC
ATG M	TTC	990 9		TAT Y
90 ATC I	144 CAC H	198 CGG R	252 ATC I	306 CAG Q
GTG V	ACA T	CTG	CTC L	ACC
GCG	ACT	ATG M	AAA K	CTC
81 CGA R	135 ATC I	189 AAC N	243 GGG	297 AAT N
CTG L	CGC R	GAC D	CAA Q	AAA K
CTG	TCG	CGG R	GAC D	CTG L
72 CGG R	126 TCG S	180 CTC L	234 ATT I	288 GAG E
GCG	GTG V	CTG	TTC	CAT H
TCC	ACC T	GAC ( D	GCT A	CTT L
63 GCG A	117 GGC	171 GGG G	225 AAG K	279 GCC A
G	AAG K	171 AGC GGG S G	GCC	CTG L
ATG M	9 9	TCC	TTA L	CGG R

## FIGURE 1A

378	432	486	540	594	648
ATC	ACT	CCT	GAG	ACA	GGA
I	T	P	E	T	G
CAG	CTT	AAC N	CGT R	CAA Q	TCC
TAT Y	CGC R	TTC	CAG Q	GAC D	TTC
369	423	477	531	585	639
GCT	CAA	GAA	ATT	GAA	ACA
A	Q	E	I	E	T
AGA	AAA	ATT	CTC	TAT	GAA
R	K	I		Y	E
GAT	ATT I	AAC N	CCT	GCT :	CTG
360	414	468	522	576	630
CTA	GTC	TAT	GAG	AAG	GTG
L	V	Y	E	K	V
GCC	GAG E	GTC V		CTA L	G
GAA	TTT	CGA	ACT	AGA	AAA
E	F	R	T	R	K
351	405	459	513	567	621
GCA	CCC	GGC	CTG	AAG	AAA
A	P	G	L	K	K
CAG	GTG V	AGT	GAC D	ATC I	CAG Q
CCA	AAT N	GCC	GAT D	GTT V	TAC Y
342	396	450	504	558	612
CTT	CTG	CCC	ATT	ACG	TAT
L	L	P	I	T	Y
ACA	AAC	CAT	9	GAG	GNA
	N	H	9	E	X
AGG	ATT I	ATT I	GTG V	CCA P	CTG
333	387	441	495	549	603
CCA	GTG	TGG	ACT	AAA	GTC
P	V	W	T	K	V
TTT	ACA	CGC	AAA	GAT	NCA
F	T	R		D	X
GGT	GAC	GCT	CCC	GAT	AAG NCA
G	D	A	P	D	K X

## FIGURE 1B

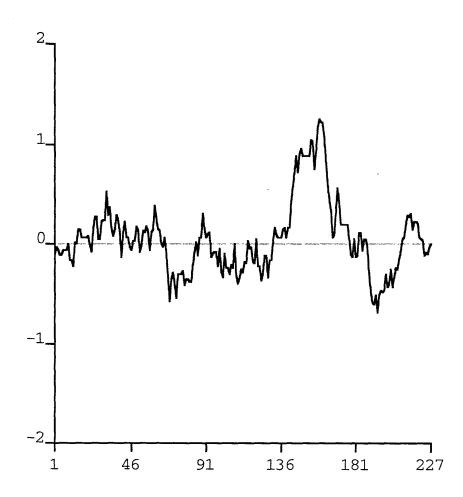
702 GTT V	756 ACT	810 TTC	
AAA K	TGT GTA	CTT	د
ACT 7	TGT	CTG	TTA
693 CAA Q	747 ATG	801 AAG	855 CTT
TTC CTA F L	738 . TGA GGA GAA A	TAG	TTA
TTC	GGA	ATT	ATA
684 GCT A	738 TGA	792 TGC	846 ATT
GTA TAT (	CCA	CCT	AAA
GTA V	ACT T	AGT	TTG
675 TAT Y	729 GTT ACT ( V T I	783 CCT 2	837 TCT
CCC	lCA	CCT	AAT
TGG W	720 AAA GCT 7 K A S	774 GGC AAA (	ATG
666 ATT I	720 AAA K	774 GGC	828 TGT
AAG K	CAG Q	ATG	GTA
AAC Z	AGC	AAG	CTA
657 ACC T	711 CCA CAA AGA P Q R	765 AGT	819 CTT
GAA E	CAA Q	AAT	CTA AGA
ACA T	CCA P	ATT	CTA

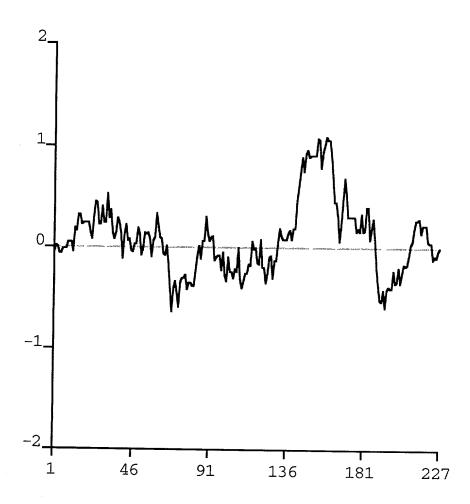
FIGURE 1C

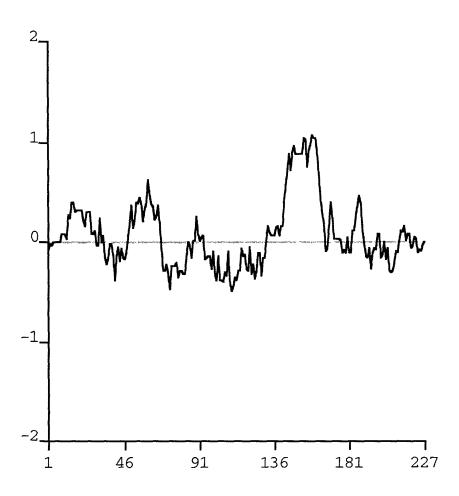
## FIGURE 2A

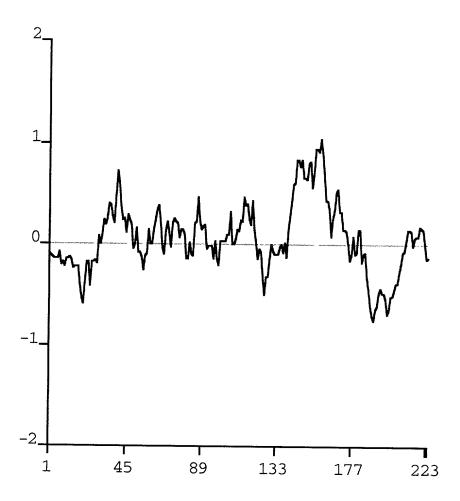
E P L I Q HWAK E P L V Q g217576 E P L I Q g450312 E P L V Q g28577	F S G T E HWAK F S G T E 9217576 F S G T E 9450312 F S G T E 928577	HMAK 9217576 9450312 928577
I K Q R L T A R W I H P A S G R V Y N I E F N P P K T V G I D D L T G E I K Q R L T A R W I H P G S G R V Y N I E F N P P K T M G I D D L T G E I K L R L T A R W I H P A S G R V Y N I E F N P P K T V G I D D L T G E L K D R L S R R W I H P P S G R V Y N L D F N P P H V H G I D D V T G E	1 1	TNKIWPYVYAFLOTKVPORSQKASVTP TNKIWPHVYAFLOTKLPORSQETSVTP TNKIRPHVYSFLOMKVPETIOKASVTP TNKIWPYVYTLFSNKITPIOSKEAY
121 121 121 121 119	161 161 161 159	201 201 201 201 199

## FIGURE 2B









Library	Lib Description	Abun	Pct Abun
MUSCNOT01		2	0.0890
COLNNOT07	colon, 60 M, match to COLNTUT16	⊣	0.0409
SEMVNOT01	_	$\boldsymbol{\vdash}$	0.0272
COLNTUT16		⊣	0.0256
SKINBIT01.	skin, leg, erythema nodosum	$\leftarrow$	0.0256
OVARTUT04	ovarian tumor, 53 F	$\leftarrow$	0.0252
ADRENOT08	adrenal gland, 20 M ·	_	0.0249
LIVSFEM03	liver/spleen, fetal M, NORM, WM	$\leftarrow$	0.0246
HIPOAZT01	brain, hippocampus, Alzheimer's, 74 M	Н	0.0245
TMLR2DT01	lymphocytes (non-adher PBMNC), M/F, 24-hr MLR	$\leftarrow$	0.0212
SYNORAB01	synovium, hip, rheumatoid, 68 F	⊣	0.0195
HNT2RAT01	hNT2 cell line, teratocarcinoma, treated RA	$\leftarrow$	0.0188
BRSTNOT07	breast, 43 F	$\leftarrow$	0.0146
PLACNOM02	placenta, neonatal F, NORM, WM	2	0.0111
NGANNOT01	ganglioneuroma, 9 M	₩	0.0073
LIVSFEM02	liver/spleen, fetal M, NORM, WM	$\vdash$	0.0026
Electronic Norther	Electronic Northern Results returned a total of 16 row(s).		

FIGURE 4